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F10920M shows a representative N-terminal restricted response to the peptide DAEFRHDSGY (SEQ ID NO:9) which covers amino acids 1-10 of the AN1792 peptide which was used as immunizing antigen.

The paragraph beginning at page 8, line 5, has been amended as follows:

B2

Fig. 20: Epitope Map: Non-restricted N-terminal response. Day 175 serum from cynomolgus monkeys was tested by ELISA against a series of 10-mer overlapping peptides (SEQ ID NOS:1-41) covering the complete AN1792 sequence. Animal number F10975F shows a representative non-restricted N-terminal response. Reactivity is seen against the two peptides N-terminal and one peptide C-terminal to the peptide DAEFRHDSGY (SEQ ID NO:9) which covers amino acids 1-10 of the AN1792 peptide.

The paragraph beginning at page 14, line 13, has been amended as follows:

B3

H₂N-Asp-Ala-Glu-Phe-Arg-His-Asp-Ser-Gly-Tyr-Glu-Val-His-His-Gln-Lys-Leu-Val-Phe-Phe-Ala-Glu-Asp-Val-Gly-Ser-Asn-Lys-Gly-Ala-Ile-Ile-Gly-Leu-Met-Val-Gly-Gly-Val-Val-Ile-Ala-OH (SEQ ID NO:42).

The paragraph beginning at page 28, line 14, has been amended as follows:

B4

Some agents for inducing an immune response contain the appropriate epitope for inducing an immune response against amyloid deposits but are too small to be immunogenic. In this situation, a peptide immunogen can be linked to a suitable carrier to help elicit an immune response. Suitable carriers include serum albumins, keyhole limpet hemocyanin, immunoglobulin molecules, thyroglobulin, ovalbumin, tetanus toxoid, or a toxoid from other pathogenic bacteria, such as diphtheria, *E. coli*, cholera, or *H. pylori*, or an attenuated toxin derivative. Other carriers include T-cell epitopes that bind to multiple MHC alleles, e.g., at least 75% of all human MHC alleles. Such carriers are sometimes known in the art as "universal T-cell epitopes." Examples of universal T-cell epitopes include:

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Influenza Hemagglutinin: HA₃₀₇₋₃₁₉ **PKYVKQNTLKLAT** (SEQ ID NO:43)

*B of
CPL*
PADRE (common residues bolded) **AKXVAAWTLKAAA** (SEQ ID NO:44)

Malaria CS: T3 epitope **EKKIAKMEKASSVFNV** (SEQ ID NO:45)

Hepatitis B surface antigen: HBsAg₁₉₋₂₈ **FFLLTRILTI** (SEQ ID NO:46)

Heat Shock Protein 65: hsp65₁₅₁₋₁₇₁ **DQSIGDLIAEAMDKGNEG** (SEQ ID NO:47)

bacille Calmette-Guerin **QVHFQPLPPAVVKL** (SEQ ID NO:48)

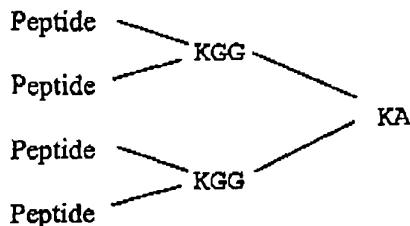
Tetanus toxoid: TT₈₃₀₋₈₄₄ **QYIKANSKFIGITEL** (SEQ ID NO:49)

Tetanus toxoid: TT₉₄₇₋₉₆₇ **FNNFTVFSFWLRVPKVSASHLE** (SEQ ID NO:50)

HIV gp120 T1: **KQIINMWQEVGKAMYA** (SEQ ID NO:51).

The paragraph beginning at page 30, line 24, has been amended as follows:

b5
The MAP4 configuration is shown below, where branched structures are produced by initiating peptide synthesis at both the N terminal and side chain amines of lysine. Depending upon the number of times lysine is incorporated into the sequence and allowed to branch, the resulting structure will present multiple N termini. In this example, four identical N termini have been produced on the branched lysine-containing core. Such multiplicity greatly enhances the responsiveness of cognate B cells.



AN90549 (A β 1-7/Tetanus toxoid 830-844 in a MAP4 configuration):

DAEFRHHDQYIKANSKFIGITEL (SEQ ID NO:52)

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AN90550 (A β 1-7/Tetanus toxoid 947-967 in a MAP4 configuration):

DAEFRHDFNNFTVSFWLRVPKVSASHLE (SEQ ID NO:53)

AN90542 (A β 1-7/Tetanus toxoid 830-844 + 947-967 in a linear configuration):

DAEFRHDQYIKANSKFIGITELFNNFTVSFWLRVPKVSASHLE

(SEQ ID NO:54)

AN90576: (A β 3-9)/Tetanus toxoid 830-844 in a MAP4 configuration):

EFRHDSGQYIKANSKFIGITEL (SEQ ID NO:55)

Peptide described in US 5,736,142 (all in linear configurations):

AN90562 (A β 1-7/ peptide) AKXVAAWTLKAAA**DAEFRHD** (SEQ ID NO:56)

AN90543 (A β 1-7 x 3/ peptide): DAEFRHDDAEFRHDDAEFRHD**AKXVAAWTLKAAA**
(SEQ ID NO:57)

Other examples of fusion proteins (immunogenic epitope of A β bolded) include
AKXVAAWTLKAAA-DAEFRHD-DAEFRHD-DAEFRHD

(SEQ ID NO:58)

DAEFRHD-AKXVAAWTLKAAA (SEQ ID NO:59)

DAEFRHD-ISQAVHAAHAEINEAGR (SEQ ID NO:60)

FRHDSGY-ISQAVHAAHABINEAGR (SEQ ID NO:61)

EFRHDSCG-ISQAVHAAHAEINEAGR (SEQ ID NO:62)

PKYVKQNTLKLAT-DAEFRHD-DAEFRHD-DAEFRHD

(SEQ ID NO:63)

DAEFRHD-PKYVKQNTLKLAT-DAEFRHD (SEQ ID NO:64)

DAEFRHD-DAEFRHD-DAEFRHD-PKYVKQNTLKLAT

(SEQ ID NO:65)

DAEFRHD-DAEFRHD-PKYVKQNTLKLAT (SEQ ID NO:66)

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Cont'd*
DAEFRHD-PKYVKQNTLKLAT-EKKIAKMEKASSVFNV-
QYIKANSKFIGITEL-FNNFTVSFWLRVPKVSASHLE-DAEFRHD
(SEQ ID NO:67)

DAEFRHD-DAEFRHD-DAEFRHD-QYIKANSKFIGITEL-
FNNFTVSFWLRVPKVSASHLE (SEQ ID NO:68)

DAEFRHD-QYIKANSKFIGITELCFNNFTVSFWLRVPKVSASHLE
(SEQ ID NO:69)

DAEFRHD-QYIKANSKFIGITELCFNNFTVSFWLRVPKVSASHLE-
DAEFRHD (SEQ ID NO:70)

DAEFRHD-QYIKANSKFIGITEL on a 2 branched resin
(SEQ ID NO:77)

peptide >
peptide > Lys-Gly-Cys

EQVTNVGGAISQAVHAAHAEINEAGR (SEQ ID NO:71) (Synuclein
fusion protein in MAP-4 configuration).

The paragraph beginning at page 60, line 24, has been amended as follows:

Preparation of coupled A β peptides: four human A β peptide conjugates (amino acid residues 1-5, 1-12, 13-28, and 33-42, each conjugated to sheep anti-mouse IgG) were prepared by coupling through an artificial cysteine added to the A β peptide using the crosslinking reagent sulfo-EMCS. The A β peptide derivatives were synthesized with the following final amino acid sequences. In each case, the location of the inserted cysteine residue is indicated by underlining. The A β 13-28 peptide derivative also had two glycine residues added prior to the carboxyl terminal cysteine as indicated.

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A β 1-12 peptide	NH2-DAEFRHDSGYEV <u>C</u> -COOH (SEQ ID NO:72)
A β 1-5 peptide	NH2-DAE <u>FRC</u> -COOH (SEQ ID NO:73)
A β 33-42 peptide	NH2- <u>C</u> -amino-heptanoic acid-GLMVGGVVIA-COOH (SEQ ID NO:74)
A β 13-28 peptide	Ac-NH-HHQKLVFFAEDVGSNK <u>GGC</u> -COOH (SEQ ID NO:75)

The paragraph beginning at page 102, line 8, has been amended as follows:

b6

The exact array of linear peptides recognized by the antibodies in the serum samples from animals immunized with AN1792 was determined by an ELISA that measured the binding of these antibodies to overlapping peptides that covered the entire A β 1-42 sequence. Biotinylated peptides with partial sequences of AN1792 were obtained from Chiron Technologies as 10 amino acid peptides with an overlap of 9 residues and a step of one residue per peptide (synthesis No. 5366, No. 5331 and No. 5814). The first 32 peptides (from the eight amino acid position upstream of the N-terminal of AN1792 down to the twenty-fourth amino acid of AN1792) are biotinylated on the C-terminal with a linker of GGK. The last 10 peptides (repeating the thirty-second peptide from the previous series) are biotinylated on the N-terminal with a linker consisting of EGEG (SEQ ID NO:76). The lyophilized biotinylated peptides were dissolved at a concentration of 5 mM in DMSO. These peptide stocks were diluted to 5 μ M in TTBS (0.05% Tween 20, 25 mM Tris HCl, 137 mM NaCl, 5.1 mM KCl, pH=7.5). 100 μ l aliquots of this 5 μ M solution were added in duplicate to streptavidin pre-coated 96-well plates (Pierce). Plates were incubated for one hour at room temperature, then washed four times with TTBS. Serum samples were diluted in specimen diluent without azide to normalize titers, and 100 μ l was added per well. These plates were incubated one hour at room temperature and then washed four times with TTBS. HRP-conjugated goat anti-human antibody (Jackson ImmunoResearch) was diluted 1:10,000 in specimen diluent without azide and 100 μ l was added per well. The plates were again